

RAW SEQUENCE LISTING  
PATENT APPLICATION US/09/208,629BDATE: 06/23/2001  
TIME: 02:15:00

INPUT SET: S36563.raw

This Raw Listing contains the General  
Information Section and those Sequences  
containing ERRORS.

## SEQUENCE LISTING

1  
2  
3 (1) General Information  
4  
5 (i) APPLICANT: Coughlin, Shaun  
6 Ishihari, Hiroaki  
7 Connolly, Andrew  
8  
9 (ii) TITLE OF THE INVENTION: Protease Activated Receptor  
10 3 and Uses Thereof  
11  
12 (iii) NUMBER OF SEQUENCES: 28  
13  
14 (iv) CORRESPONDENCE ADDRESS:  
15 (A) ADDRESSEE: MORRISON & FOERSTER  
16 (B) STREET: 755 Page Mill Road  
17 (C) CITY: Palo Alto  
18 (D) STATE: CA  
19 (E) COUNTRY: USA  
20 (F) ZIP: 94304-1018  
21  
22 (v) COMPUTER READABLE FORM:  
23 (A) MEDIUM TYPE: Diskette  
24 (B) COMPUTER: IBM Compatible  
25 (C) OPERATING SYSTEM: DOS  
26 (D) SOFTWARE: FastSEQ for Windows Version 2.0  
27  
28 (vi) CURRENT APPLICATION DATA:  
29 (A) APPLICATION NUMBER: 09/208,629  
30 (B) FILING DATE: 08-DEC-1998  
31 (C) CLASSIFICATION:  
32  
33 (vii) PRIOR APPLICATION DATA:  
34 (A) APPLICATION NUMBER: 08/742,440  
35 (B) FILING DATE: 30-OCT-1996  
36  
37  
38  
39 (viii) ATTORNEY/AGENT INFORMATION:  
40 (A) NAME: Catherine M. Polizzi  
41 (B) REGISTRATION NUMBER: 40,130  
42 (C) REFERENCE/DOCKET NUMBER: 220002060310  
43  
44 (ix) TELECOMMUNICATION INFORMATION:  
45 (A) TELEPHONE: (650) 813-5600

Does Not Comply  
Corrected Diskette Needed

# RAW SEQUENCE LISTING PATENT APPLICATION US/09/208,629B

DATE: 06/23/2001  
TIME: 02:15:01

INPUT SET: S36563.raw

46 (B) TELEFAX: (650) 494-0792  
47 (C) TELEX:  
48  
49

## ERRORED SEQUENCES FOLLOW:

123 (2) INFORMATION FOR SEQ ID NO:3:  
124  
125 (i) SEQUENCE CHARACTERISTICS:  
--> 126 (A) LENGTH: 407 amino acids 4 shown  
127 (B) TYPE: amino acid  
128 (C) STRANDEDNESS: single  
129 (D) TOPOLOGY: linear  
130  
131 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:  
132  
133 Thr Leu Tyr Thr  
134

201 (2) INFORMATION FOR SEQ ID NO:6:  
202  
203 (i) SEQUENCE CHARACTERISTICS:  
--> 204 (A) LENGTH: 408 amino acids 8 shown  
205 (B) TYPE: amino acid  
206 (C) STRANDEDNESS: single  
207 (D) TOPOLOGY: linear  
208  
209 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:  
210  
211 Cys Ser Met Ile Leu Gln Ile Ser  
212 1 5 ← number the amino acids under every 5

595 (2) INFORMATION FOR SEQ ID NO:24: amino acids  
596  
597 (i) SEQUENCE CHARACTERISTICS:  
598 (A) LENGTH: 380 amino acids  
599 (B) TYPE: amino acid  
600 (C) STRANDEDNESS: single  
601 (D) TOPOLOGY: linear  
602 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:24: ← add mandatory heading  
603 Gln His Pro Val Ala Gly Ser Gln Asp Ile Lys Met Lys Ile Leu Ile  
--> 604 1 5 10 15  
--> 605 Leu Val Ala Ala Gly Leu Leu Phe Leu Pro Val Thr Val Cys Gln Ser  
--> 606 20 25 30  
--> 607 Gly Ile Asn Val Ser Asp Asn Ser Ala Lys Pro Thr Leu Thr Ile Lys  
--> 608 35 40 45  
--> 609 Ser Phe Asn Gly Gly Pro Gln Asn Thr Phe Glu Glu Phe Pro Leu Ser  
--> 610 50 55 60  
--> 611 Asp Ile Glu Gly Trp Thr Gly Ala Thr Thr Thr Ile Lys Ala Glu Cys

# RAW SEQUENCE LISTING PATENT APPLICATION US/09/208,629B

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```

--> 612 65 70 75 80
--> 613 Pro Glu Asp Ser Ile Ser Thr Leu His Val Asn Asn Ala Thr Ile Gly
--> 614 85 90 95
--> 615 Tyr Leu Arg Ser Ser Leu Ser Thr Gln Val Ile Pro Ala Ile Tyr Ile
--> 616 100 105 110
--> 617 Leu Leu Phe Val Val Gly Val Pro Ser Asn Ile Val Thr Leu Trp Lys
--> 618 115 120 125
--> 619 Leu Ser Leu Arg Thr Lys Ser Ile Ser Leu Val Ile Phe His Thr Asn
--> 620 130 135 140
--> 621 Leu Ala Ile Ala Asp Leu Leu Phe Cys Val Thr Leu Pro Phe Lys Ile
--> 622 145 150 155 160
--> 623 Ala Tyr His Leu Asn Gly Asn Asn Trp Val Phe Gly Glu Val Met Cys
--> 624 165 170 175
--> 625 Arg Ile Thr Thr Val Val Phe Tyr Gly Asn Met Tyr Cys Ala Ile Leu
--> 626 180 185 190
--> 627 Ile Leu Thr Cys Met Gly Ile Asn Arg Tyr Leu Ala Thr Ala His Pro
--> 628 195 200 205
--> 629 Phe Thr Tyr Gln Lys Leu Pro Lys Arg Ser Phe Ser Leu Leu Met Cys
--> 630 210 215 220
--> 631 Gly Ile Val Trp Val Met Val Phe Leu Tyr Met Leu Pro Phe Val Ile
--> 632 225 230 235 240
--> 633 Leu Lys Gln Glu Tyr His Leu Val His Ser Glu Ile Thr Thr Cys His
--> 634 245 250 255
--> 635 Asp Val Val Asp Ala Cys Glu Ser Pro Ser Ser Phe Arg Phe Tyr Tyr
--> 636 260 265 270
--> 637 Phe Val Ser Leu Ala Phe Phe Gly Phe Leu Ile Pro Phe Val Ile Ile
--> 638 275 280 285
--> 639 Ile Phe Cys Tyr Thr Thr Leu Ile His Lys Leu Lys Ser Lys Asp Arg
--> 640 290 295 300
--> 641 Ile Trp Leu Gly Tyr Ile Lys Ala Val Leu Leu Ile Leu Val Ile Phe
--> 642 305 310 315 320
--> 643 Thr Ile Cys Phe Ala Pro Thr Asn Ile Ile Leu Val Ile His His Ala
--> 644 325 330 335
--> 645 Asn Tyr Tyr Tyr His Asn Thr Asp Ser Leu Tyr Phe Met Tyr Leu Ile
--> 646 340 345 350
--> 647 Ala Leu Cys Leu Gly Ser Leu Asn Ser Cys Leu Asp Pro Phe Leu Tyr
--> 648 355 360 365
--> 649 Phe Val Met Ser Lys Val Val Asp Gln Leu Asn Pro
--> 650 370 375 380
--> 651

```

OK

These  
are in  
bold  
print  
due to  
error  
on  
previous  
page

652 (2) INFORMATION FOR SEQ ID NO:25:

653

654 (i) SEQUENCE CHARACTERISTICS:

655 (A) LENGTH: 7 amino acids

656 (B) TYPE: amino acid

657 (C) STRANDEDNESS: single

658 (D) TOPOLOGY: linear

659

660 Ser Ala Met Ala Arg Pro Leu

--&gt; 661 1 5

662

← insert heading

663

(2) INFORMATION FOR SEQ ID NO:26:

RAW SEQUENCE LISTING  
PATENT APPLICATION US/09/208,629BDATE: 06/23/2001  
TIME: 02:15:02

INPUT SET: S36563.raw

664  
665 (i) SEQUENCE CHARACTERISTICS:  
666 (A) LENGTH: 13 amino acids  
667 (B) TYPE: amino acid  
668 (C) STRANDEDNESS: single  
669 (D) TOPOLOGY: linear  
670

*insert leading*

--> 671 Arg Pro Arg Arg Asp Ile Trp Glu Asp Ile His Ala Trp  
672 1 5 10  
673

674 (2) INFORMATION FOR SEQ ID NO:27:  
675

676 (i) SEQUENCE CHARACTERISTICS:  
677 (A) LENGTH: 393 amino acids  
678 (B) TYPE: amino acid  
679 (C) STRANDEDNESS: single  
680 (D) TOPOLOGY: linear  
681

*insert leading*

--> 682 Cys Ser Met Ile Leu Gln Ile Ser Xaa Arg Leu Arg Asp Gly Thr Gln  
683 1 5 10 15  
--> 684 Val Ile Lys Met Lys Ala Leu Ile Phe Ala Ala Gly Leu Leu  
685 20 25 30  
--> 686 Leu Leu Pro Thr Phe Cys Gln Ser Gly Met Glu Asn Asp Thr Asn Asn  
687 35 40 45  
--> 688 Leu Ala Lys Pro Thr Leu Pro Ile Lys Thr Phe Arg Gly Ala Pro Pro  
689 50 55 60  
--> 690 Asn Ser Phe Glu Glu Phe Pro Phe Ser Ala Leu Glu Gly Trp Thr Gly  
691 65 70 75 80  
--> 692 Ala Thr Ile Thr Val Lys Ile Lys Cys Pro Glu Glu Ser Ala Ser His  
693 85 90 95  
--> 694 Leu His Val Lys Asn Ala Thr Met Gly Tyr Leu Thr Ser Ser Leu Ser  
695 100 105 110  
--> 696 Thr Lys Leu Ile Pro Ala Ile Tyr Leu Leu Val Phe Val Val Gly Val  
697 115 120 125  
--> 698 Pro Ala Asn Ala Val Thr Leu Trp Met Leu Phe Phe Arg Thr Arg Ser  
699 130 135 140  
--> 700 Ile Cys Thr Thr Val Phe Tyr Thr Asn Leu Ala Ile Ala Asp Phe Leu  
701 145 150 155 160  
--> 702 Phe Cys Val Thr Leu Pro Phe Lys Ile Ala Tyr His Leu Asn Gly Asn  
703 165 170 175  
--> 704 Asn Trp Val Phe Gly Glu Val Leu Cys Arg Ala Thr Thr Val Ile Phe  
705 180 185 190  
--> 706 Tyr Gly Asn Met Tyr Cys Ser Ile Leu Leu Leu Ala Cys Ile Ser Ile  
707 195 200 205  
--> 708 Asn Arg Tyr Leu Ala Ile Val His Pro Phe Thr Tyr Arg Gly Leu Pro  
709 210 215 220  
--> 710 Lys His Thr Tyr Ala Leu Val Thr Cys Gly Leu Val Trp Ala Thr Val  
711 225 230 235 240  
--> 712 Phe Leu Tyr Met Leu Pro Phe Phe Ile Leu Lys Gln Glu Tyr Tyr Leu  
713 245 250 255  
--> 714 Val Gln Pro Asp Ile Thr Thr Cys His Asp Val His Asn Thr Cys Glu  
715 260 265 270

*O/C*

# RAW SEQUENCE LISTING PATENT APPLICATION US/09/208,629B

DATE: 06/23/2001  
TIME: 02:15:02

INPUT SET: S36563.raw

```

--> 716 Ser Ser Ser Pro Phe Gln Leu Tyr Tyr Phe Ile Ser Leu Ala Phe Phe
--> 717      275      280      285
--> 718 Gly Phe Leu Ile Pro Phe Val Leu Ile Ile Tyr Cys Tyr Ala Ala Ile
--> 719      290      295      300
--> 720 Ile Arg Thr Leu Asn Ala Tyr Asp His Arg Trp Leu Trp Tyr Val Lys
--> 721      305      310      315      320
--> 722 Ala Ser Leu Leu Ile Leu Val Ile Phe Thr Ile Cys Phe Ala Pro Ser
--> 723      325      330      335
--> 724 Asn Ile Ile Leu Ile Ile His His Ala Asn Tyr Tyr Tyr Asn Asn Thr
--> 725      340      345      350
--> 726 Asp Gly Leu Tyr Phe Ile Tyr Leu Ile Ala Leu Cys Leu Gly Ser Leu
--> 727      355      360      365
--> 728 Asn Ser Cys Leu Asp Pro Phe Leu Tyr Phe Leu Met Ser Lys Thr Arg
--> 729      370      375      380
--> 730 Asn His Ser Thr Ala Tyr Leu Thr Lys
--> 731      385      390
--> 732

```

O/C

733 (2) INFORMATION FOR SEQ ID NO:28:

734

735 (i) SEQUENCE CHARACTERISTICS:

736 (A) LENGTH: 13 amino acids

737 (B) TYPE: amino acid

738 (C) STRANDEDNESS: single

739 (D) TOPOLOGY: linear

740

741 Asn Asp Leu Arg Glu Gln Gly Gln Pro Ser Gln Arg Thr

--> 742 1 5 10

743

744

# SEQUENCE VERIFICATION REPORT

## PATENT APPLICATION US/09/208,629B

DATE: 06/23/2001  
TIME: 02:15:02

INPUT SET: S36563.raw

| Line | Error  | Original Text   |
|------|--|---|
| 126  | Entered (407) and Calc. Seq. Length (4) differ | (A) LENGTH: 407 amino acids                                 |
| 204  | Entered (408) and Calc. Seq. Length (8) differ | (A) LENGTH: 408 amino acids                                 |
| 604  | Response Exceeds Line Limitations              | 1 5 10 15   |
| 605  | Response Exceeds Line Limitations              | Leu Val Ala Ala Gly Leu Leu Phe Leu Pro Val Thr Val Cys     |
| 606  | Response Exceeds Line Limitations              | 20 25 30  |
| 607  | Response Exceeds Line Limitations              | Gly Ile Asn Val Ser Asp Asn Ser Ala Lys Pro Thr Leu Thr I   |
| 608  | Response Exceeds Line Limitations              | 35 40 45  |
| 609  | Response Exceeds Line Limitations              | Ser Phe Asn Gly Gly Pro Gln Asn Thr Phe Glu Glu Phe Pro     |
| 610  | Response Exceeds Line Limitations              | 50 55 60  |
| 611  | Response Exceeds Line Limitations              | Asp Ile Glu Gly Trp Thr Gly Ala Thr Thr Thr Ile Lys Ala G   |
| 612  | Response Exceeds Line Limitations              | 65 70 75  |
| 613  | Response Exceeds Line Limitations              | Pro Glu Asp Ser Ile Ser Thr Leu His Val Asn Asn Ala Thr I   |
| 614  | Response Exceeds Line Limitations              | 85 90 95  |
| 615  | Response Exceeds Line Limitations              | Tyr Leu Arg Ser Ser Leu Ser Thr Gln Val Ile Pro Ala Ile Ty  |
| 616  | Response Exceeds Line Limitations              | 100 105 110   |
| 617  | Response Exceeds Line Limitations              | Leu Leu Phe Val Val Gly Val Pro Ser Asn Ile Val Thr Leu T   |
| 618  | Response Exceeds Line Limitations              | 115 120 125   |
| 619  | Response Exceeds Line Limitations              | Leu Ser Leu Arg Thr Lys Ser Ile Ser Leu Val Ile Phe His Th  |
| 620  | Response Exceeds Line Limitations              | 130 135 140   |
| 621  | Response Exceeds Line Limitations              | Leu Ala Ile Ala Asp Leu Leu Phe Cys Val Thr Leu Pro Phe     |
| 622  | Response Exceeds Line Limitations              | 145 150 155   |
| 623  | Response Exceeds Line Limitations              | Ala Tyr His Leu Asn Gly Asn Asn Trp Val Phe Gly Glu Va      |
| 624  | Response Exceeds Line Limitations              | 165 170 175   |
| 625  | Response Exceeds Line Limitations              | Arg Ile Thr Thr Val Val Phe Tyr Gly Asn Met Tyr Cys Ala     |
| 626  | Response Exceeds Line Limitations              | 180 185 190   |
| 627  | Response Exceeds Line Limitations              | Ile Leu Thr Cys Met Gly Ile Asn Arg Tyr Leu Ala Thr Ala     |
| 628  | Response Exceeds Line Limitations              | 195 200 205   |
| 629  | Response Exceeds Line Limitations              | Phe Thr Tyr Gln Lys Leu Pro Lys Arg Ser Phe Ser Leu Leu     |
| 630  | Response Exceeds Line Limitations              | 210 215 220   |
| 631  | Response Exceeds Line Limitations              | Gly Ile Val Trp Val Met Val Phe Leu Tyr Met Leu Pro Phe     |
| 632  | Response Exceeds Line Limitations              | 225 230 235   |
| 633  | Response Exceeds Line Limitations              | Leu Lys Gln Glu Tyr His Leu Val His Ser Glu Ile Thr Thr C   |
| 634  | Response Exceeds Line Limitations              | 245 250 255   |
| 635  | Response Exceeds Line Limitations              | Asp Val Val Asp Ala Cys Glu Ser Pro Ser Ser Phe Arg Phe     |
| 636  | Response Exceeds Line Limitations              | 260 265 270   |
| 637  | Response Exceeds Line Limitations              | Phe Val Ser Leu Ala Phe Phe Gly Phe Leu Ile Pro Phe Val I   |
| 638  | Response Exceeds Line Limitations              | 275 280 285   |
| 639  | Response Exceeds Line Limitations              | Ile Phe Cys Tyr Thr Thr Leu Ile His Lys Leu Lys Ser Lys A   |
| 640  | Response Exceeds Line Limitations              | 290 295 300   |
| 641  | Response Exceeds Line Limitations              | Ile Trp Leu Gly Tyr Ile Lys Ala Val Leu Leu Ile Leu Val Ile |
| 642  | Response Exceeds Line Limitations              | 305 310 315   |
| 643  | Response Exceeds Line Limitations              | Thr Ile Cys Phe Ala Pro Thr Asn Ile Ile Leu Val Ile His His |
| 644  | Response Exceeds Line Limitations              | 325 330 335   |
| 645  | Response Exceeds Line Limitations              | Asn Tyr Tyr Tyr His Asn Thr Asp Ser Leu Tyr Phe Met Ty      |
| 646  | Response Exceeds Line Limitations              | 340 345 350   |
| 647  | Response Exceeds Line Limitations              | Ala Leu Cys Leu Gly Ser Leu Asn Ser Cys Leu Asp Pro Phe     |
| 648  | Response Exceeds Line Limitations              | 355 360 365   |
| 649  | Response Exceeds Line Limitations              | Phe Val Met Ser Lys Val Val Asp Gln Leu Asn Pro             |

# SEQUENCE VERIFICATION REPORT

## PATENT APPLICATION US/09/208,629B

DATE: 06/23/2001  
TIME: 02:15:03

INPUT SET: S36563.raw

| Line | Error                             | Original Text  |
|------|-----------------------------------|--|
| 650  | Response Exceeds Line Limitations | 370 375 380  |
| 661  | Response Exceeds Line Limitations | 1 5  |
| 672  | Response Exceeds Line Limitations | 1 5 10   |
| 683  | Response Exceeds Line Limitations | 1 5 10 15  |
| 684  | Response Exceeds Line Limitations | Val Ile Lys Met Lys Ala Leu Ile Phe Ala Ala Ala Gly Leu L  |
| 685  | Response Exceeds Line Limitations | 20 25 30   |
| 686  | Response Exceeds Line Limitations | Leu Leu Pro Thr Phe Cys Gln Ser Gly Met Glu Asn Asp Th     |
| 687  | Response Exceeds Line Limitations | 35 40 45   |
| 688  | Response Exceeds Line Limitations | Leu Ala Lys Pro Thr Leu Pro Ile Lys Thr Phe Arg Gly Ala    |
| 689  | Response Exceeds Line Limitations | 50 55 60   |
| 690  | Response Exceeds Line Limitations | Asn Ser Phe Glu Glu Phe Pro Phe Ser Ala Leu Glu Gly Trp    |
| 691  | Response Exceeds Line Limitations | 65 70 75   |
| 692  | Response Exceeds Line Limitations | Ala Thr Ile Thr Val Lys Ile Lys Cys Pro Glu Glu Ser Ala S  |
| 693  | Response Exceeds Line Limitations | 85 90 95   |
| 694  | Response Exceeds Line Limitations | Leu His Val Lys Asn Ala Thr Met Gly Tyr Leu Thr Ser Ser    |
| 695  | Response Exceeds Line Limitations | 100 105 110  |
| 696  | Response Exceeds Line Limitations | Thr Lys Leu Ile Pro Ala Ile Tyr Leu Leu Val Phe Val Val G  |
| 697  | Response Exceeds Line Limitations | 115 120 125  |
| 698  | Response Exceeds Line Limitations | Pro Ala Asn Ala Val Thr Leu Trp Met Leu Phe Phe Arg Th     |
| 699  | Response Exceeds Line Limitations | 130 135 140  |
| 700  | Response Exceeds Line Limitations | Ile Cys Thr Thr Val Phe Tyr Thr Asn Leu Ala Ile Ala Asp P  |
| 701  | Response Exceeds Line Limitations | 145 150 155  |
| 702  | Response Exceeds Line Limitations | Phe Cys Val Thr Leu Pro Phe Lys Ile Ala Tyr His Leu Asn    |
| 703  | Response Exceeds Line Limitations | 165 170 175  |
| 704  | Response Exceeds Line Limitations | Asn Trp Val Phe Gly Glu Val Leu Cys Arg Ala Thr Thr Va     |
| 705  | Response Exceeds Line Limitations | 180 185 190  |
| 706  | Response Exceeds Line Limitations | Tyr Gly Asn Met Tyr Cys Ser Ile Leu Leu Ala Cys Ile S      |
| 707  | Response Exceeds Line Limitations | 195 200 205  |
| 708  | Response Exceeds Line Limitations | Asn Arg Tyr Leu Ala Ile Val His Pro Phe Thr Tyr Arg Gly    |
| 709  | Response Exceeds Line Limitations | 210 215 220  |
| 710  | Response Exceeds Line Limitations | Lys His Thr Tyr Ala Leu Val Thr Cys Gly Leu Val Trp Ala    |
| 711  | Response Exceeds Line Limitations | 225 230 235  |
| 712  | Response Exceeds Line Limitations | Phe Leu Tyr Met Leu Pro Phe Phe Ile Leu Lys Gln Glu Tyr    |
| 713  | Response Exceeds Line Limitations | 245 250 255  |
| 714  | Response Exceeds Line Limitations | Val Gln Pro Asp Ile Thr Thr Cys His Asp Val His Asn Thr    |
| 715  | Response Exceeds Line Limitations | 260 265 270  |
| 716  | Response Exceeds Line Limitations | Ser Ser Ser Pro Phe Gln Leu Tyr Tyr Phe Ile Ser Leu Ala P  |
| 717  | Response Exceeds Line Limitations | 275 280 285  |
| 718  | Response Exceeds Line Limitations | Gly Phe Leu Ile Pro Phe Val Leu Ile Ile Tyr Cys Tyr Ala Al |
| 719  | Response Exceeds Line Limitations | 290 295 300  |
| 720  | Response Exceeds Line Limitations | Ile Arg Thr Leu Asn Ala Tyr Asp His Arg Trp Leu Trp Tyr    |
| 721  | Response Exceeds Line Limitations | 305 310 315  |
| 722  | Response Exceeds Line Limitations | Ala Ser Leu Leu Ile Leu Val Ile Phe Thr Ile Cys Phe Ala Pr |
| 723  | Response Exceeds Line Limitations | 325 330 335  |
| 724  | Response Exceeds Line Limitations | Asn Ile Ile Leu Ile Ile His His Ala Asn Tyr Tyr Tyr Asn As |
| 725  | Response Exceeds Line Limitations | 340 345 350  |
| 726  | Response Exceeds Line Limitations | Asp Gly Leu Tyr Phe Ile Tyr Leu Ile Ala Leu Cys Leu Gly    |
| 727  | Response Exceeds Line Limitations | 355 360 365  |

**SEQUENCE VERIFICATION REPORT**  
**PATENT APPLICATION US/09/208,629B**DATE: 06/23/2001  
TIME: 02:15:03**INPUT SET: S36563.raw**

| Line | Error                             | Original Text  |
|------|-----------------------------------|--|
| 728  | Response Exceeds Line Limitations | Asn Ser Cys Leu Asp Pro Phe Leu Tyr Phe Leu Met Ser Ly |
| 729  | Response Exceeds Line Limitations | 370                      375                      380  |
| 730  | Response Exceeds Line Limitations | Asn His Ser Thr Ala Tyr Leu Thr Lys                    |
| 731  | Response Exceeds Line Limitations | 385                      390                           |
| 742  | Response Exceeds Line Limitations | 1                      5                      10       |